

T. neapolitana DLE (2779) SOM

Page N 48

October 13, 1994 (Thursday)

I infected DH12S cells with the ϕ from #5, #6, #7 (2ml cells grown in 2x YT + 5 μ l ϕ ; grown 37°C (air shaker) 16 hrs)

RF isolated by alkaline/SOS except 1 μ l RNase A (1mg/ml) added to prep at NH₄OH addition

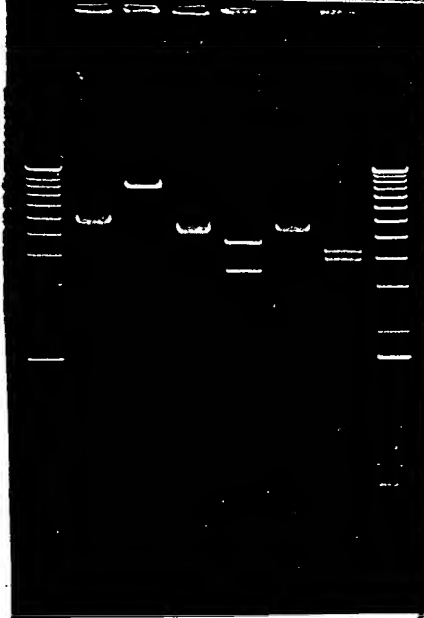
NA dissolved in 50 μ l T₁₀E₁

DIGEST SCHEME

(React 3)	HOP	10 μ l	✓	Incubated 37°C (heat block) 1 hour.
	10x Bfr	2	✓	
	DNA	7	✓	
(4 μ l)	Eco47III	1	✓	
	Total	20 μ l		

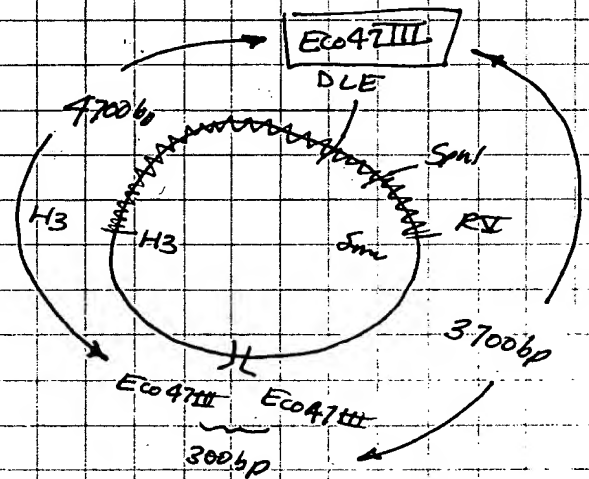
3% Agarose Gel (1xTAE); 190V/1h

1. Run #2779 Eco47III DyeS



6 Kb
5 Kb
4 Kb
3 Kb
2 Kb

Comment:



The bands are migrating at the expected distances for #6. There must have been an overabundance of some "component" causing the DNA to run faster. I will clone into the pUC vector.

To Page No. 50

ss d & Understo d by me,

Date

Invented by

Date

May forgo

10/24/94

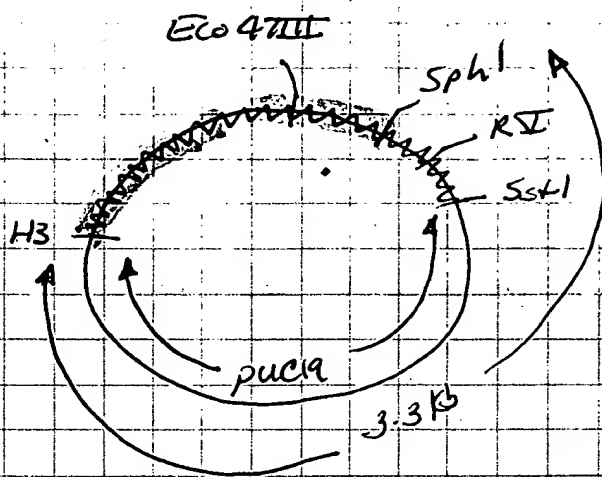
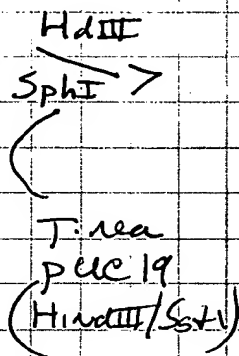
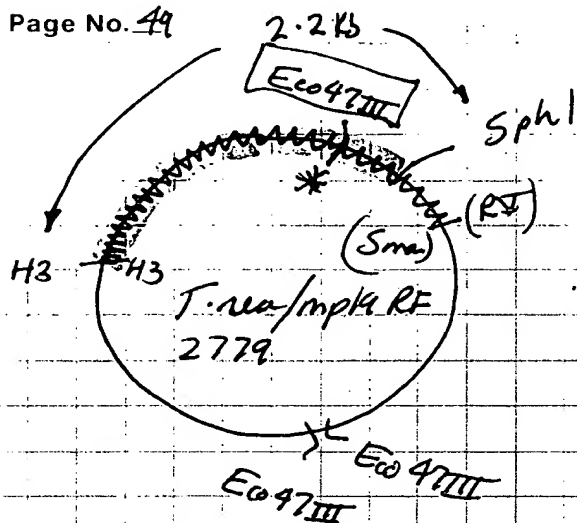
Record d by

Pr j ct No. 20222

10-13-94

From Page No. 49

October 13, 1994 (Thurs)



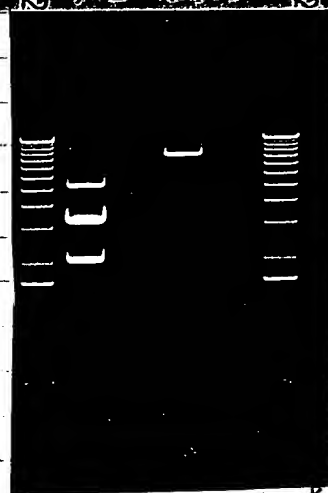
DIGEST SCHEME

		<i>T. nea/puc</i>	#6	<i>2779/mp19</i>		
(React 2)	HOM	12 μ e	✓	6 μ e	✓	Incubated 37°C (heat)
	10X Bfr	2	✓	2	✓	
	DNA	4	✓	12	✓	
(104 μ e)	HindIII	1	✓	1	✓	
(104 μ e)	SphI	1	✓	1	✓	
	Torac	20 μ e		20 μ e		1:10 → 1:50

0.8% Agarose Gel (1XTAE), 190



Comments: I should see a 3.3 Kb (desired fragment) and a 2.2 Kb fragment from the *T. nea/puc19* clone and I do. Unfortunately I should see a 7.9 Kb fragment and 2.2 Kb (desired fragment) from the *T. nea/mp19* (2779 #6) RF DNA and I don't. Both sites were present before I performed the mutagenesis (see p. 35) - I will have to repeat.



To Page No

Witnessed & Understood by me,

Date

Invented by

Date

M. Jones

10/24/94

Recorded by

D. J. Schmitt

10-13-94

T. neapolitana SDM

ge N _____

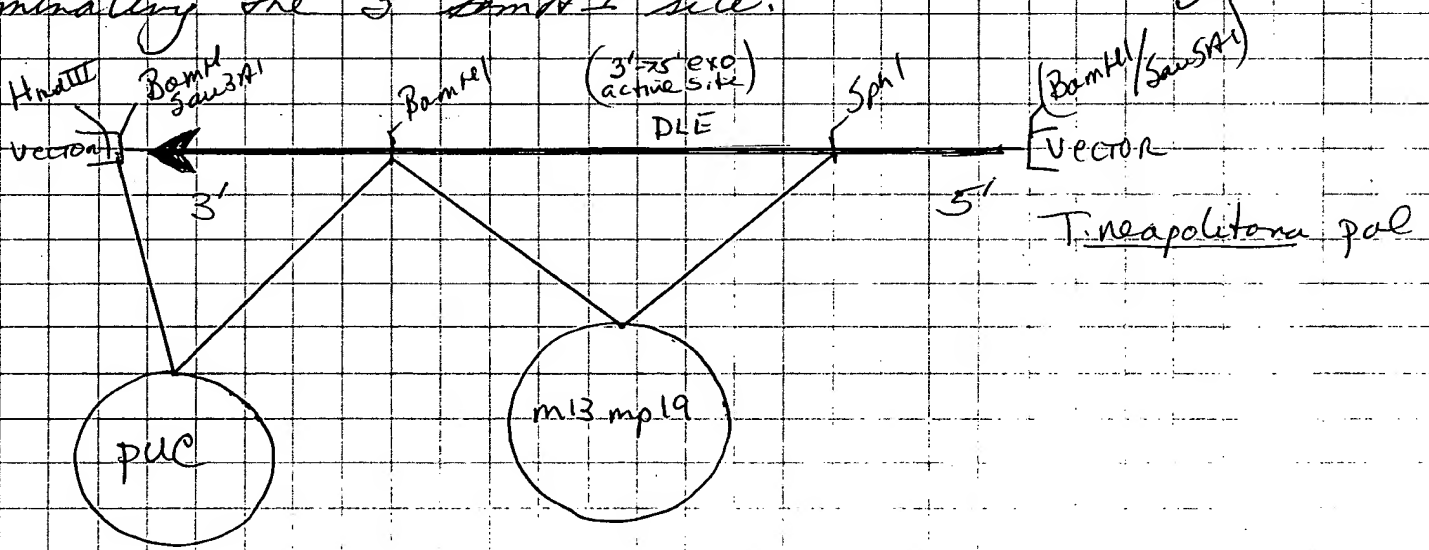
January 25, 1995 (Wednesday)

I'm BAAACK!!

after a tour of duty with Joel's group, a trip to Aulba and a few week vacation. If I am back and ready to administer the fatal blow to this project. I will finish sequencing this gene, mutagenize it to conform to our needs, and overexpress it so people can enough enzyme to swim in it and still have money left over for a cup of coffee and a copy of the New York Times!

How's that for an opening!

First things first. Let's reclone the region of the *pal* gene we are interested in mutagenizing. Rob and I have had no success with the last clone. Secondly, let's make the subclone more user friendly by eliminating the 3' BamHI site.



stryg BamHI/Sau3A
 PCR
 Clone Hd3/BamHI

make ssDNA
 D → A by SDM

To Page No. 52

ed & Understood by me, <i>Ray Jorgo</i>	Date 1/27/95 4/24/95	Inv nt d by <i>Ray Jorgo</i>	Date 1-25-95
Recorded by <i>Ray Jorgo</i>			

From Page No. 51

January 25, 1995 (Wednes.)

T. neapolitana pSPORT DNA made by Michael Smith
(not the Nobel Laureate; the Horebag)

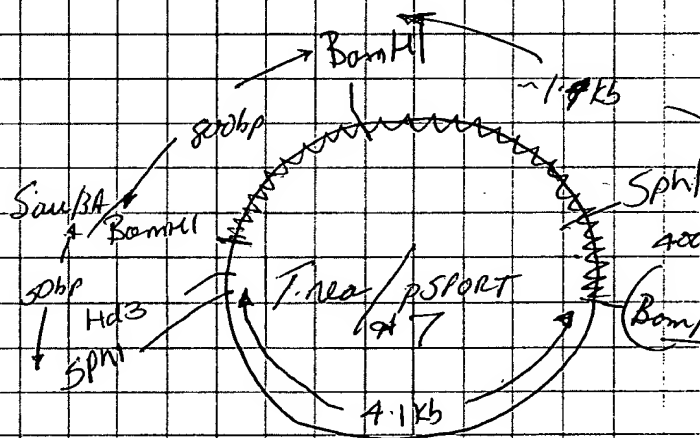
DIGEST SCHEME

	<u>T. nea/pSPORT</u>	<u>mBmp19</u>	<u>pUC18</u>
(React 6) HOH	15 μ l ✓	13 μ l ✓	13 μ l ✓
10X B/R	2 ✓	2 ✓	2 ✓
DNA	1 ✓	3 ✓	3 ✓
(10 μ l) BamHI	1 ✓	1 ✓	1 ✓
(10 μ l) SphI	1 ✓	1 ✓	1 ✓
Form	20 μ l	20 μ l	20 μ l
(0.14 μ l) CAP			1 μ l

Incubated 37°C (heat-block) 1:00 \rightarrow 2:45

0.8% Agarose Gel (1XTAE)
190 Volts

BamHI/SphI
12/27/95



I forgot to run the 1Kb ladder.
what a hore bag!

Fragment T. nea/pSPORT 7 BamHI/SphI shows
the 4.5 Kb, 1.4 Kb, 0.8 Kb, 0.25 Kb.
Perhaps something partialled. Try again
but do it separately.

Witnessed & Understood by me,

Date

Invnt d by

Date

May Jones

1/27/95

R cord d by

Dr. Michael Smith

1-25-95

T. neapolitana SDM

Project N 20222
B k N 3884

Exhibit S-3
Appl. No. 09/558,421

Tag N 52

January 26, 1995 (Thursday)

DIGEST SCHEME:

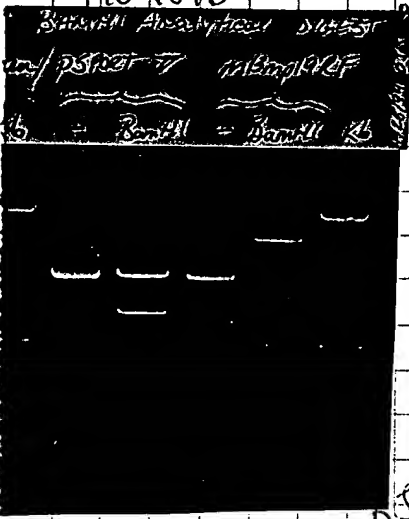
		<i>T. nea</i> /pSPORT		m13 mp19 (~270ng/μl)	
1 (React 3)	HOH	24.5 μl	✓	22.5 μl	✓
	10X B/R	3	✓	3	✓
	DNA	1	✓	3	✓
F107 (104/μl)	BamHI	1.5	✓	1.5	✓
	TOTAL	30 μl		30 μl	

Inoculated 37°C (heat block) 8:04 → 9:08

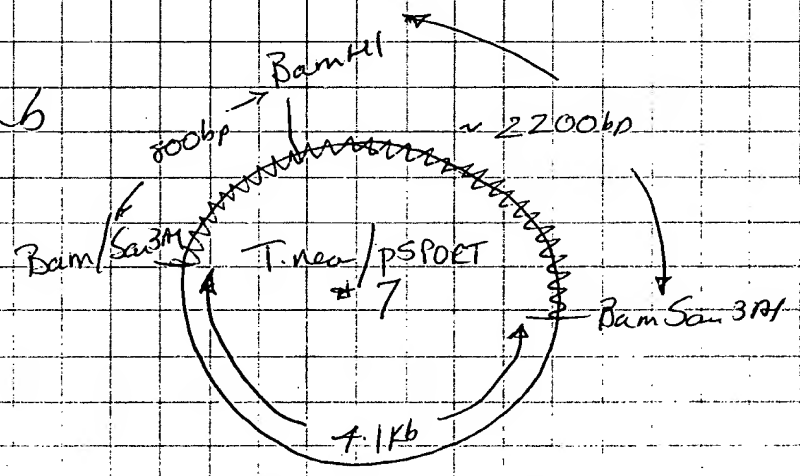
✓ 3 μl removed for analytical gel.

		<i>T. nea</i> /pSPORT	mp19	
DIGEST	27 μl	✓	✓	Inoculated 37°C (heat block) 9:17 → 10:25
1M KCl	2	✓	✓	
HOH	9	✓	✓	
1/4 μl SphI	2	✓	✓	
TOTAL	40 μl			

Agarose Gel (1XTBE)
190 Volts

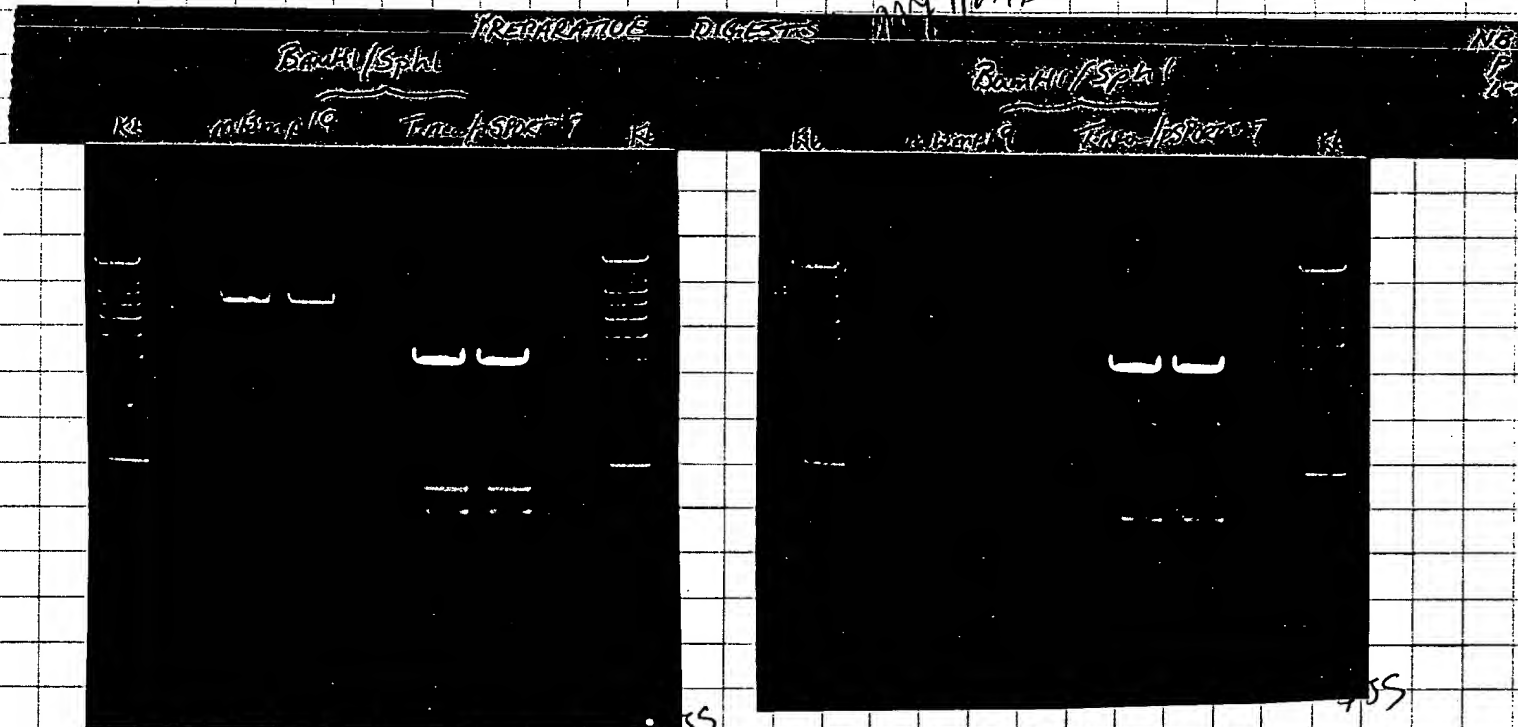


Comments



To Page No. _____

is d & Understood by me,	Date	Invent d by	Dat
Mer Torgo	1/27/95	Drumf. Schmidt	1-26-95

From Page No. 53January 26, 1995 (th0.8% Agarose Gel (1X TAE); Run at 190 Volts1/27/95

Bands extracted from the gel and placed in the same tube. The DNA was purified away from the agarose using Gene Clean as described by the manufacturer (BIO-101)

DNA eluted in 14 μ l H₂O

LIGATION SCHEME

ET6402 (Ligase)	DNA	14 μ l	✓
5X Bfr		4	✓
1 μ l (Ligase)		2	✓
Form		20 μ l	

Incubated 22°C (room-temp)
2'15" → 3'15"

→ 1 μ l ligation / 2 μ l for transformation

To Page

Witnessed & Understood by me,

Mary Loup

Date

1/27/95

Investigated by

Recorded by

David J. Kimmel

Date

1-26-95

T. neapolitana 50M

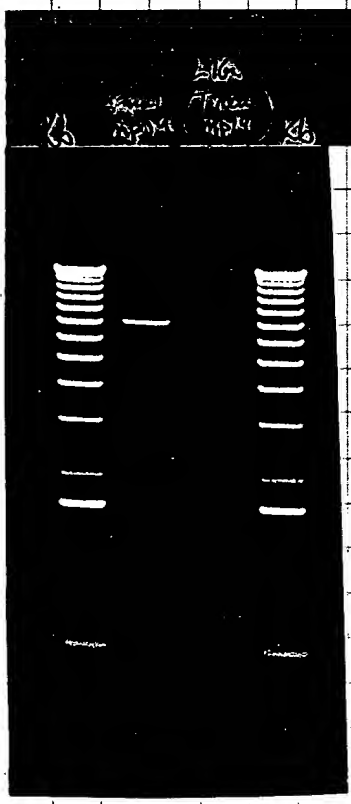
Pr j ct N .
B k N .

age No. 54

January 26, 1995 (Thursday)

DH10B Electrocompetent
20 μ l DH10B Electrocompetent Cells + 1 μ l (of a 1/3 dilution; Serp 54)
2.5 KV
1ml LB, 37°C au shaken 20 min
 ↳ 10% applied to LB plate in 4ml Soft Agar (0.7%)
 90% + IPTG (1mM) and X-gal 100 μ l of 4%

incubated 37°C incubated



1127/95

To Page No. _____

Read & Understood by me, <i>May Longo</i>	Date 1/27/95	Invent d by <i>Dr. J. Schmidt</i>	Date 1-26-95
		Recorded by	

56

Project No. 20222

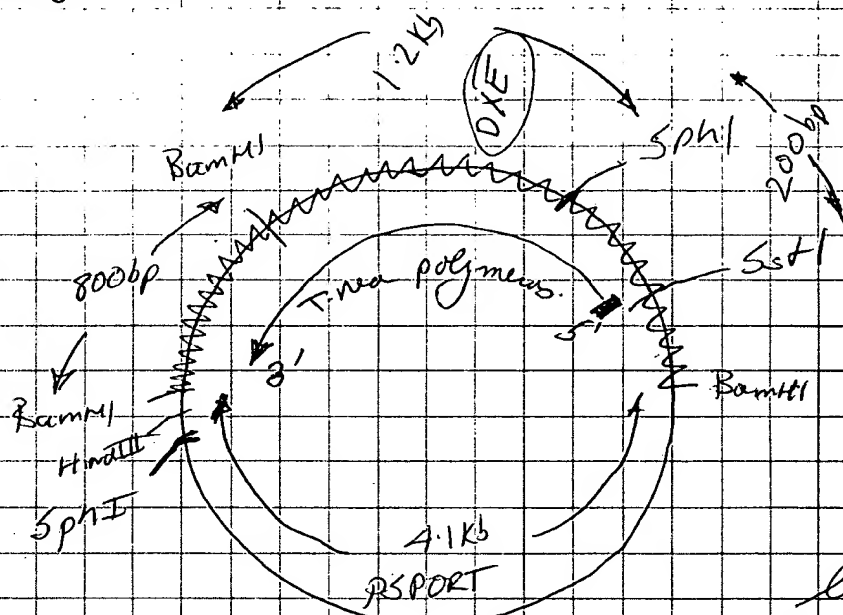
Book No. 3884

TITLE

T. neapolitana 30M

From Page N 3

February 7, 1995 (Tuesday)



I can clone a SphI fragment into m13mp19 and let m13 determine which direction is best suited.

I can subclone the fragment into an expression vector with SphI / HindIII.

DIGEST SCHEME

	T. nea pSPORT	(272 ng/μl) m13mp19	
(Ready) HOH	21 μl ✓	20 μl ✓	9:42 am →
10X Bfr	3 ✓	3 ✓	
DNA	3 ✓	3 ✓	
CMG105 (100 μl) SphI	3 ✓	3 ✓	
(0.10 μl) CAP	0 ✓	1 ✓	9:46 am →
Toraz.	30 μl	30 μl	

Run on 0.8% Agarose Gel at 75 Volts

To Page

Witnessed & Understood by me,

May Longo

Date

2/16/95

Inv. nted by

Recorded by

[Signature]

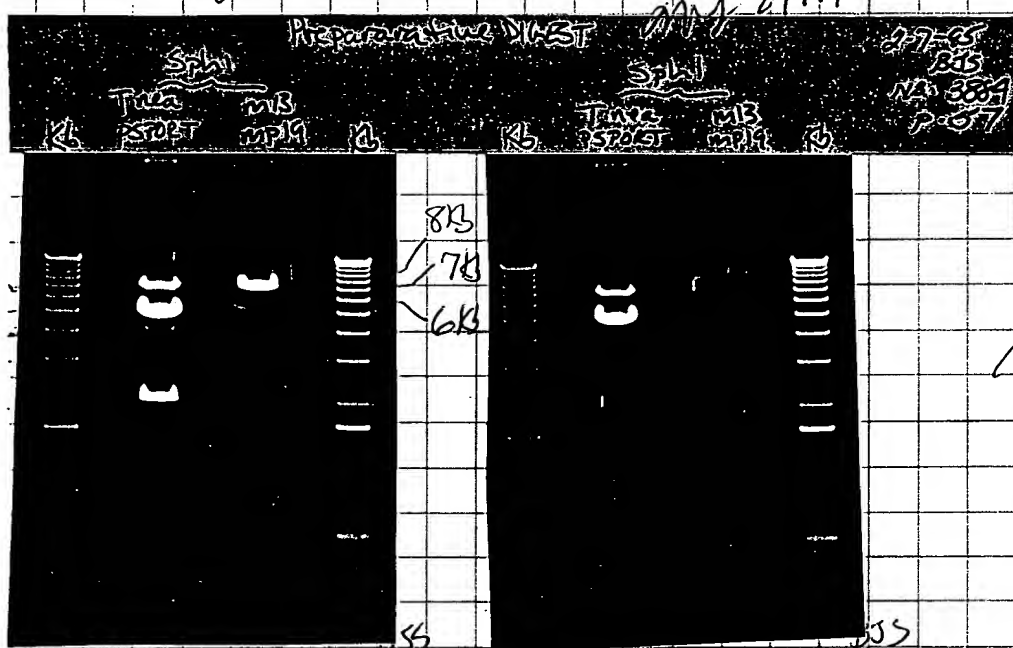
Date

2-7-95

g No. 56

February 7, 1995 (Tuesday)

0.8% Agarose Gel (1X TAE); 75 V, 16



Bands extracted from the gel and the DNA purified away from the agarose using Gene Clean as described by 8.10-10.1. DNA eluted in 14 µl H₂O.

LIGATION SCHEME

H ₂ O	-	µl	
5X Bfr	4		✓
DNA	14		✓
WAT DNA Ligase	2		✓
TOTAL	20	µl	

Incubated 3:23 pm → 4:03 at room-temperature (~22°C)
3 µl removed for transformation

152 F' IQ Competent Cell Transformation
1 µl competent cells + 3 µl ligation (see above)
2 min on ice, 35 seconds at 42°C water bath
1 and 90 µl applied to LB + No Antibiotic plates in 4 ml 0.7% Top Agar + 100 µl 2% X-Gal + 10 µl 100 mM IPTG
incubated 16 hours at 37°C incubator

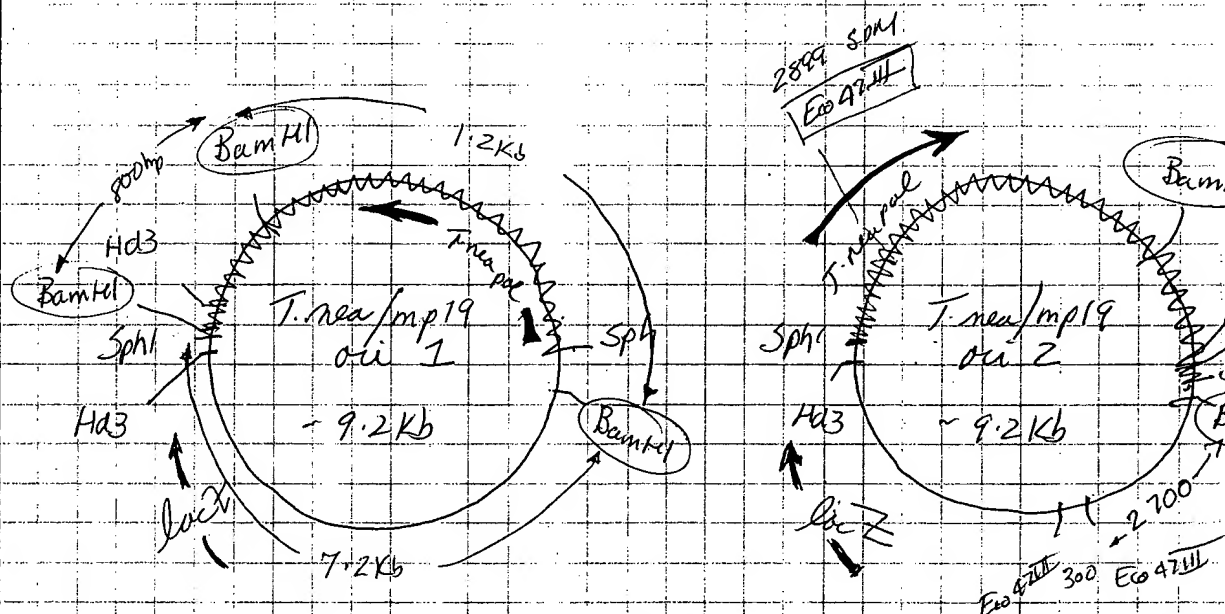
To Page No. 58

ed & Understood by me, Ney Torso	Date 2/16/95	Invented by [Signature]	Date 2-7-95
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From Page No. 57

February 8, 1985

- I added 200 μ l of DH5 α F' IQ lawn cells to 10 ml Arice Brown
- I added 1 ml of the cells to 8 glass tubes
- Each tube was inoculated with a clear plaque and incubated at 37°C (8:00 am →



SphI	7.2 Kb	VECTOR	7.2 Kb	VECTOR
	2 Kb	INSERT	2 Kb	INSERT
BamHI	7.2 Kb	VECTOR	8.4 Kb	VECTOR + I
	1.2 Kb	} INSERT	0.8 Kb	INSERT
	0.8 Kb			

Add *T. nea*/pSPORT as a positive control for both digests

To Page

Witnessed & Understood by me, M. J. J. J.	Date 2/11/85	Invented by M. J. J. J.	Date 2-8-85
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T-neapolitaner SDM

Book No. 3884

59

ag No. 58

February 8, 1995 (Wednesday)

DIGEST SCHEMES

1	PER RXN	x 9 =	COCKTAIL	
HOH	7 μ	x 9 =	63 μ	<input checked="" type="checkbox"/>
React 6) 10x Bfr	2	x 9 =	18 μ	<input checked="" type="checkbox"/>
DNA	10			
(100 μ l) SpH1	1	x 9 =	9 μ	<input checked="" type="checkbox"/>
TOTAL	20 μ		90 μ	

For T-neapolitaner
control add

TDE 20 μ ☒
DNA 2 μ ☒
TOTAL 22 μ ☒

✓

add 10 μ to reaction

	PER RXN	x 9 =	COCKTAIL	
HOH	7 μ	x 9 =	63 μ	<input checked="" type="checkbox"/>
React 3) 10x Bfr	2	x 9 =	18	<input checked="" type="checkbox"/>
DNA	10			
(100 μ l) BsmH1	1	x 9 =	9	<input checked="" type="checkbox"/>
TOTAL	20 μ		90 μ	

Continued on page 1 of Notebook 3966

Arani Patel

Gel photo

To Page No. _____

Sed & Understood by me,

Date

Invented by

Date

May Longo

2/16/95

Recorded by

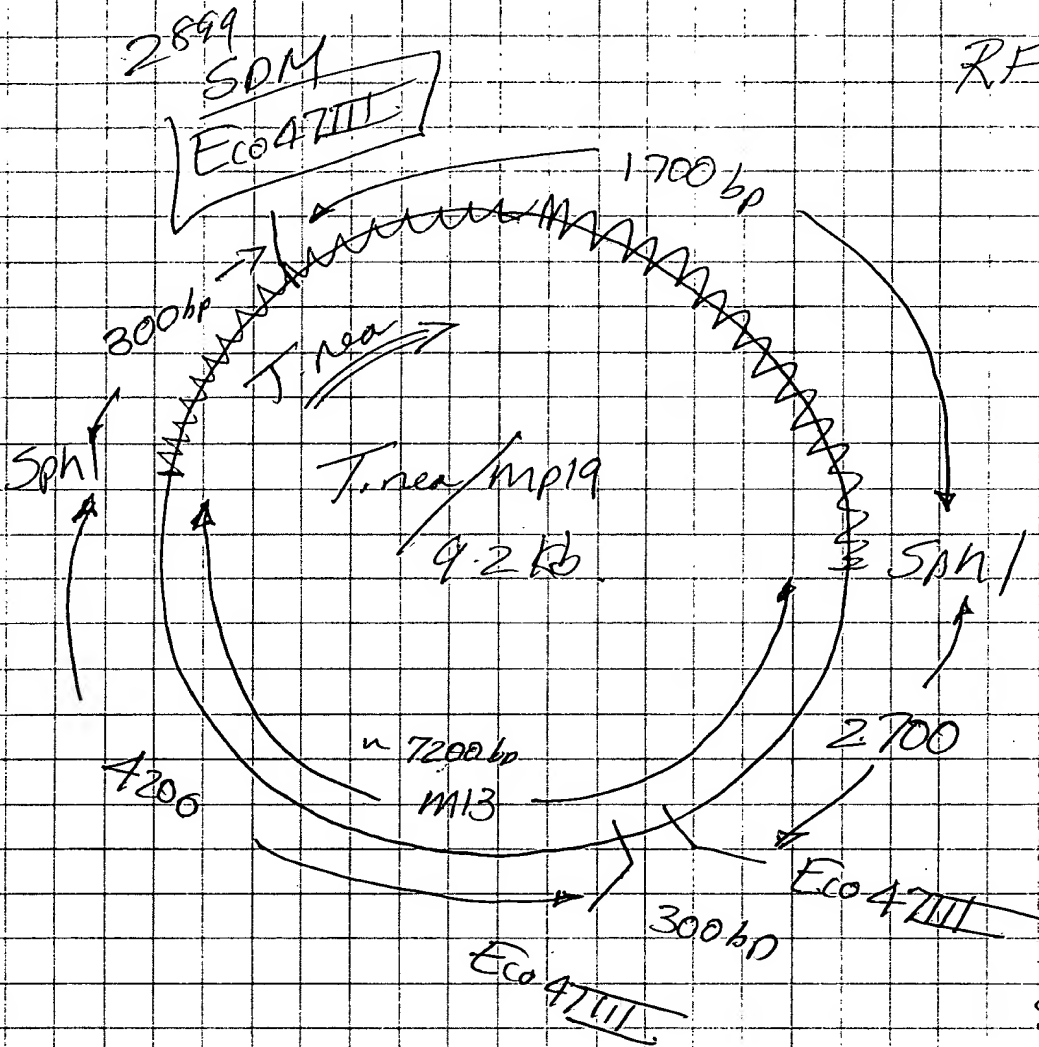
Brenda L. Lamm

2-8-95

Page No. _____

SDM 2899

RF map



See on file

	Eco 47III		PARENT	MUTANT
PARENT	8.9 Kb 0.3 Kb	5Kb 4Kb	=====	=====
MUTANT	4.5 Kb 4.4 Kb 0.3 Kb	3Kb 2Kb 1.6Kb 11Kb 500bp	=====	=====

may be too light to see

Read & Understood by me, Mary J. Info	Date 2/16/95	Invented by Dr. J. K. Smith	Date 2-16-95
	Record d by Dr. J. K. Smith		

The mutant Phe to Ala

g No. _____

① The same phenyl alanine corresponding to Tag polymerase will be changed to tyrosine

② For exo D will be changed to Alanine (corresponding region of Tag).

Brian cloned the SphI fragment of the Pol into M13mp.

I isolated the single stranded DNA from CJ236 as described before in Bio rad manual.

Test 5µl ssDNA

The DNA looks real good.

For D-A (3'-5' exo mutant oligo) is

5' GA | CGT | TTC | AAG | CGC | TAG | GGC | AAA | ABA # 2899
Eco47III site

For Phe → Tyr (O-helix)

~~GA~~ GTA | TAT | TAT | AGA | GTA | GTT | AAC | CAT | CTC | TCC | A
2904

kinased 2899 before.

kinased 2904 as follows:

2µl oligo (210 pmol)
6µl 5X buff (350mM Tris pH 7.6, 50mM MgCl₂, 50mM KCl, 5mM P.M.E.)
1µl 10mM ATP
0.5µl T4 Kinase (50)
20.5µl H₂O
5' at 37°C → heat at 65°C + 3µl TE
T Page 10/22

sed & Und rstood by m ,

H. T. J.

Date

4/8/95

Invented by

Record d by

H. T. J.

Date

3/14/95